

This Page Is Inserted by IFW Operations
and is not a part of the Official Record

BEST AVAILABLE IMAGES

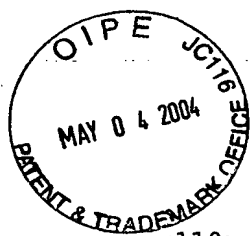
Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

**As rescanning documents *will not* correct images,
please do not report the images to the
Image Problem Mailbox.**



SEQUENCE LISTING

<110> ADLER, JON ELLIOT
ZOZULYA, SERGEY
LI, XIADONG
O'CONNELL, SHAWN
STASZEWSKI, LENA

<120> T1R TASTE RECEPTORS AND GENES ENCODING SAME

<130> 078003/0277870/RXT

<140> 09/799,629

<141> 2001-03-07

<150> 60/187,546

<151> 2000-03-07

<150> 60/195,536

<151> 2000-04-07

<150> 60/209,840

<151> 2000-06-06

<150> 60/214,213

<151> 2000-06-23

<150> 30/226,448

<151> 2000-08-17

<150> 60/259,227

<151> 2001-01-03

<160> 20

<170> PatentIn Ver. 2.1

<210> 1

<211> 876

<212> DNA

<213> Homo sapiens

<400> 1

```
agcctggcag tggcctcagg cagagtctga cgcgcacaaa ctttcaggcc caggaagcga 60
ggacaccact ggggccccag ggtgtggcaa gtgaggatgg caagggtttt gctaaacaaa 120
tcctctgccc gctccccgcc ccgggctcac tccatgtgag gcccagtcg gggcagccac 180
ctgccgtgcc tgttggaagt tgccctctgcc atgctgggcc ctgctgtcct gggcctcagc 240
ctctgggctc tcctgcaccc tgggacgggg gcccattgt gcctgtcaca gcaacttagg 300
atgaaggggg actacgtgct gggggggctg ttccccctgg gcgaggccga ggaggctggc 360
ctccgcagcc ggacacggcc cagcagccct gtgtgcacca ggtacagagg tgggacggcc 420
tgggtcgggg tcagggtgac caggtctggg gtgctcctga gctggggccg aggtggccat 480
ctgcggttct gtgtggcccc aggttctcct caaacggcct gctctgggca ctggccatga 540
aaatggccgt ggaggagatc aacaacaagt cggatctgct gcccgggctg cgcttgggct 600
acgacctctt tgatacgtgc tcggagcctg tgggtggccat gaagcccagc ctcatgttcc 660
tggccaaggc aggcagccgc gacatcgccg cctactgcaa ctacacgcag taccagcccc 720
gtgtgctggc tgtcatcggg cccactcgt cagagctcgc catggtcacc ggcaagttct 780
tcagcttctt cctcatgccc cagtggggcg cccccacca tcaccacccc ccaaccaacc 840
cctgccccgt gggagccctt tgtgtcagga gaatgc 876
```

<210> 2
 <211> 2687
 <212> DNA
 <213> Homo sapiens

<400> 2
 tacatgcacc ccaccagacc ctgccctggg agccctgtgt cagaagatgc tcttggcctt 60
 gcaggtcagc tacggtgcta gcatggagct gctgagcgcc cgggagacct tcccctcctt 120
 cttccgcacc gtgcccagcg accgtgtgca gctgacggcc gccgcggagc tgctgcagga 180
 gttcggctgg aactgggtgg ccgccctggg cagcgacgac gagtacggcc ggcagggcct 240
 gagcatcttc tcggccctgg ccgcggcacg cggcatctgc atcgcgacg agggcctggt 300
 gccgctgccc cgtgccgatg actcgcggtt ggggaagggt caggacgtcc tgcaccaggt 360
 gaaccagagc agcgtgcagg tgggtgctgt gttcgcctcc gtgcacgccg cccacgccct 420
 cttcaactac agcatcagca gcaggctctc gcccagggtg tgggtggcca gcgaggcctg 480
 gctgacctct gacctggtca tggggctgcc cggcatggcc cagatgggca cggtgcttgg 540
 cttccctcag aggggtgccc agctgcacga gttccccag tacgtgaaga cgcacctggc 600
 cctggccacc gaccggcctt tctgctctgc cctgggagag agggagcagg gtctggagga 660
 ggacgtggtg ggccagcgct gcccgagtg tgactgcac acgctgcaga acgtgagcgc 720
 agggctaaat caccaccaga cgttctctgt ctacgcagct gtgtatagcg tggcccaggc 780
 cctgcacaac actcttcagt gcaacgcctc aggttgcccc gcgcaggacc ccgtgaagcc 840
 ctggcaggtg agcccgggag atgggggtgt gctgtcctct gcatgtgccc agggccaccag 900
 gcacggccac cacgcctgag ctggagggtg ctggcggtc agccccgtcc cccgcccgc 960
 gtcctggag aacatgtaca acctgacctt ccacgtgggc gggctgccgc tgcggttcga 1020
 cagcagcgga aacgtggaca tggagtacga cctgaagctg tgggtgtggc agggctcagt 1080
 gcccaaggct cagcagctgg gcaggttcaa cggcagctc aggcagagc gcctgaagat 1140
 ccgctggcac acgtctgaca accaggtgag gtgagggtgg gtgtgccagg cgtgcccgtg 1200
 gtagcccccg cggcagggcg cagcctgggg gtggggggcg ttccagtctc ccgtgggcat 1260
 gccagccga gcagagccag accccaggcc tgtgcgcaga agcccgtgtc ccggtgctcg 1320
 cggcagtgcc agggaggcca ggtgcgccgg gtcaaggggt tccactcctg ctgctacgac 1380
 tgtgtggact gcgagggcgg cagctaccgg caaaacccag gtgagccgcc tccccggcag 1440
 gcgggggtgg gaacgcagca ggggagggtc ctgccaaagt ctgactctga gaccagagcc 1500
 cacagggtac aagacgaaca cccagcgccc ttctcctctc tcacagacga catcgctgc 1560
 accttttgtg gccaggatga gtggtccccg gacggaagca cacgtgctt ccgcgcagg 1620
 tctcggttcc tggcatgggg cgagccggct gtgctgctgc tgctcctgct gctgagcctg 1680
 gcgctggggc ttgtgctggc tgccttgggg ctgttcgttc accatcggga cagcccactg 1740
 gttcaggcct cggggggggc cctggcctgc tttggcctgg tgtgcctggg cctggtctgc 1800
 ctacgcgtcc tectgttccc tggccagccc agccctgccc gatgcctggc ccagcagccc 1860
 ttgtcccacc tcccgtcac gggctgcctg agcacactct tectgcaggc ggccgagatc 1920
 ttctgtgagt cagaactgcc tctgagctgg gcagaccggc tgagtggctg cctgcggggg 1980
 cctgggcctt ggctgggtgt gctgctggcc atgctgggtg aggtcgcaact gtgcacctgg 2040
 tacctggtgg ccttcccggc ggaggtgggt acggaactgg acatgctgcc cacggaggcg 2100
 ctggtgcact gccgcacacg ctctgggtc agcttcggcc tagcgcacgc caccaatgcc 2160
 acgctggcct ttctctgctt cctgggcact ttctgggtgc ggagccagcc gggctgctac 2220
 aacctggccc gtggcctcac ctttgccatg ctggcctact tcatcacctg ggtctccttt 2280
 gtgcccctcc tggccaatgt gcaggtggtc ctcaggcccc ccgtgcagat gggcgccctc 2340
 ctgctctgtg tectgggcat cctggctgcc ttccacctgc ccaggtgtta cctgctcatg 2400
 cggcagccag ggtcaaacac ccccagttc ttctgggag gggggcctgg ggatgcccaa 2460
 ggccagaatg acgggaacac aggaaatcag gggaaacatg agtgacccaa ccctgtgatc 2520
 tcagccccgg tgaaccagca cttagctgcg atccccccca agccagcaat gaccctgtgc 2580
 tcgctacaga gaccctcccg ctctagggtt tgaccccagg ttgtctcctg accctgaccc 2640
 cacagtgage cctaggcctg gagcacgtgg acaccctgt gaccatc 2687

<210> 3
 <211> 2553

<212> DNA

<213> Homo sapiens

<400> 3

```

atgctggggc ctgctgtcct gggcctcagc ctctgggctc tectgcaccc tgggacgggg 60
gccccattgt gcctgtcaca gcaacttagg atgaaggggg actacgtgct ggggggggctg 120
ttccccctgg gcgaggccga ggaggctggc ctccgcagcc ggacacggcc cagcagccct 180
gtgtgcacca ggttctcttc aaacggcctg ctctgggcac tggccatgaa aatggccgtg 240
gaggagatca acaacaagtc ggatctgctg cccgggctgc gcctgggcta cgacctcttt 300
gatacgtgct cggagcctgt ggtggccatg aagcccagcc tcatgttcct ggccaaggca 360
ggcagccgcg acatcgccgc ctactgcaac tacacgcagt accagccccg tgtgtctggct 420
gtcatcgggc cccactcgtc agagctcgcc atggtcaccg gcaagttctt cagcttcttc 480
ctcatgcccc actacgggtg tagcatggag ctgctgagcg cccgggagac cttccccctcc 540
ttcttccgca ccgtgcccag cgaccgtgtg cagctgacgg ccgcccggga gctgctgcag 600
gagttcggct ggaactgggt ggccgcccctg ggcagcgacg acgagtacgg ccggcagggc 660
ctgagcatct tctcgccctt ggccgcgga cgcggcatct gcacgcgca cgaggccctg 720
gtgccgctgc cccgtgcga tgactcgcg ctggggaagg tgcaggacgt cctgcaccag 780
gtgaaccaga gcagcgtgca ggtggtgctg ctgttcgcct ccgtgcacgc cgcccacgcc 840
ctcttcaact acagcatcag cagcaggctc tcgcccagg tgtgggtggc cagcagggcc 900
tggctgacct ctgacctggt catggggctg cccggcatgg cccagatggg cacggtgctt 960
ggcttctctc agaggggtgc ccagctgcac gagttcccc agtacgtgaa gacgcacctg 1020
gccctggcca ccgaccggc cttctgctct gccctgggcg agagggagca gggctctggag 1080
gaggacgtgg tgggcccagc ctgcccgcag tgtgactgca tcacgctgca gaacgtgagc 1140
gcagggctaa atcaccacca gacgttctct gtctacgcag ctgtgtatag cgtggcccag 1200
gcctgcaca acactcttca gtgcaacgcc tcaggctgce ccgcgcagga ccccgtaag 1260
ccctggcagc tectggagaa catgtacaac ctgaccttcc acgtgggagg gctgcgctg 1320
cggttcgaca gcagcggaaa cgtggacatg gagtacgacc tgaagctgtg ggtgtggcag 1380
ggctcagtgc ccaggtcca cgacgtgggc aggttcaacg gcagcctcag gacagagcgc 1440
ctgaagatcc gctggcacac gtctgacaac cagaagcccg tgtcccgggtg ctcgcggcag 1500
tgccaggagg gccaggtgcg ccgggtcaag gggttccact cctgctgcta cgactgtgtg 1560
gactgagagg cgggcagcta ccggcaaaac ccagacgaca tcgcctgcac cttttgtggc 1620
caggatgagt ggtccccgga gcgaagcaca cgctgcttcc gccgcaggtc tcggttctctg 1680
gcatggggcg agccggtgt gctgctgctg ctctgctgce tgagcctggc gctgggcctt 1740
gtgctggctg ctttgggggt gttcgttcac catcgggaca gcccactggt tcaggcctcg 1800
ggggggcccc tggcctgctt tggcctgggt tgctgggccc tggctgcct cagcgtcttc 1860
ctgttccctg gccagcccag ccctgcccga tgccctggccc agcagccctt gtcccacctc 1920
ccgctcacgg gctgcctgag cacactcttc ctgcaggcgg ccgagatctt cgtggagtca 1980
gaactgcctc tgagctgggc agaccggctg agtggctgcc tgccggggccc ctgggcctgg 2040
ctgggtggtg tgctggccat gctggtggag gtgcactgt gcacctggta cctggtggcc 2100
ttcccgcgg aggtggtgac ggactggcac atgctgccc cggaggcgct ggtgactgc 2160
cgcacacgct cctgggtcag cttcggccta gcgcacgcca ccaatgccac gctggccttt 2220
ctctgcttcc tgggcacttt cctggtgceg agccagccgg gctgctacaa ccgtgcccgt 2280
ggcctcacct ttgccatgct ggcctacttc atcacctggg tctcctttgt gcccctcctg 2340
gccaatgtgc aggtggtcct caggcccgcc gtgcagatgg gcgccctcct gctctgtgtc 2400
ctgggcatcc tggctgcctt ccacctgccc aggtgttacc tgctcatgcg gcagccaggg 2460
ctcaacaccc ccgagttctt cctgggaggg ggccctgggg atgcccagg ccagaatgac 2520
gggaacacag gaaatcaggg gaaacatgag tga 2553

```

<210> 4

<211> 850

<212> PRT

<213> Homo sapiens

<400> 4

```

Met Leu Gly Pro Ala Val Leu Gly Leu Ser Leu Trp Ala Leu Leu His
  1                      5                      10                      15

```

Pro Gly Thr Gly Ala Pro Leu Cys Leu Ser Gln Gln Leu Arg Met Lys
 20 25 30

Gly Asp Tyr Val Leu Gly Gly Leu Phe Pro Leu Gly Glu Ala Glu Glu
 35 40 45

Ala Gly Leu Arg Ser Arg Thr Arg Pro Ser Ser Pro Val Cys Thr Arg
 50 55 60

Phe Ser Ser Asn Gly Leu Leu Trp Ala Leu Ala Met Lys Met Ala Val
 65 70 75 80

Glu Glu Ile Asn Asn Lys Ser Asp Leu Leu Pro Gly Leu Arg Leu Gly
 85 90 95

Tyr Asp Leu Phe Asp Thr Cys Ser Glu Pro Val Val Ala Met Lys Pro
 100 105 110

Ser Leu Met Phe Leu Ala Lys Ala Gly Ser Arg Asp Ile Ala Ala Tyr
 115 120 125

Cys Asn Tyr Thr Gln Tyr Gln Pro Arg Val Leu Ala Val Ile Gly Pro
 130 135 140

His Ser Ser Glu Leu Ala Met Val Thr Gly Lys Phe Phe Ser Phe Phe
 145 150 155 160

Leu Met Pro His Tyr Gly Ala Ser Met Glu Leu Leu Ser Ala Arg Glu
 165 170 175

Thr Phe Pro Ser Phe Phe Arg Thr Val Pro Ser Asp Arg Val Gln Leu
 180 185 190

Thr Ala Ala Ala Glu Leu Leu Gln Glu Phe Gly Trp Asn Trp Val Ala
 195 200 205

Ala Leu Gly Ser Asp Asp Glu Tyr Gly Arg Gln Gly Leu Ser Ile Phe
 210 215 220

Ser Ala Leu Ala Ala Ala Arg Gly Ile Cys Ile Ala His Glu Gly Leu
 225 230 235 240

Val Pro Leu Pro Arg Ala Asp Asp Ser Arg Leu Gly Lys Val Gln Asp
 245 250 255

Val Leu His Gln Val Asn Gln Ser Ser Val Gln Val Val Leu Leu Phe
 260 265 270

Ala Ser Val His Ala Ala His Ala Leu Phe Asn Tyr Ser Ile Ser Ser
 275 280 285

Arg Leu Ser Pro Lys Val Trp Val Ala Ser Glu Ala Trp Leu Thr Ser
 290 295 300

Asp Leu Val Met Gly Leu Pro Gly Met Ala Gln Met Gly Thr Val Leu
 305 310 315 320

Gly Phe Leu Gln Arg Gly Ala Gln Leu His Glu Phe Pro Gln Tyr Val
 325 330 335
 Lys Thr His Leu Ala Leu Ala Thr Asp Pro Ala Phe Cys Ser Ala Leu
 340 345 350
 Gly Glu Arg Glu Gln Gly Leu Glu Glu Asp Val Val Gly Gln Arg Cys
 355 360 365
 Pro Gln Cys Asp Cys Ile Thr Leu Gln Asn Val Ser Ala Gly Leu Asn
 370 375 380
 His His Gln Thr Phe Ser Val Tyr Ala Ala Val Tyr Ser Val Ala Gln
 385 390 395 400
 Ala Leu His Asn Thr Leu Gln Cys Asn Ala Ser Gly Cys Pro Ala Gln
 405 410 415
 Asp Pro Val Lys Pro Trp Gln Leu Leu Glu Asn Met Tyr Asn Leu Thr
 420 425 430
 Phe His Val Gly Gly Leu Pro Leu Arg Phe Asp Ser Ser Gly Asn Val
 435 440 445
 Asp Met Glu Tyr Asp Leu Lys Leu Trp Val Trp Gln Gly Ser Val Pro
 450 455 460
 Arg Leu His Asp Val Gly Arg Phe Asn Gly Ser Leu Arg Thr Glu Arg
 465 470 475 480
 Leu Lys Ile Arg Trp His Thr Ser Asp Asn Gln Lys Pro Val Ser Arg
 485 490 495
 Cys Ser Arg Gln Cys Gln Glu Gly Gln Val Arg Arg Val Lys Gly Phe
 500 505 510
 His Ser Cys Cys Tyr Asp Cys Val Asp Cys Glu Ala Gly Ser Tyr Arg
 515 520 525
 Gln Asn Pro Asp Asp Ile Ala Cys Thr Phe Cys Gly Gln Asp Glu Trp
 530 535 540
 Ser Pro Glu Arg Ser Thr Arg Cys Phe Arg Arg Arg Ser Arg Phe Leu
 545 550 555 560
 Ala Trp Gly Glu Pro Ala Val Leu Leu Leu Leu Leu Leu Ser Leu
 565 570 575
 Ala Leu Gly Leu Val Leu Ala Ala Leu Gly Leu Phe Val His His Arg
 580 585 590
 Asp Ser Pro Leu Val Gln Ala Ser Gly Gly Pro Leu Ala Cys Phe Gly
 595 600 605
 Leu Val Cys Leu Gly Leu Val Cys Leu Ser Val Leu Leu Phe Pro Gly
 610 615 620

Gln	Pro	Ser	Pro	Ala	Arg	Cys	Leu	Ala	Gln	Gln	Pro	Leu	Ser	His	Leu	625	630	635	640
Pro	Leu	Thr	Gly	Cys	Leu	Ser	Thr	Leu	Phe	Leu	Gln	Ala	Ala	Glu	Ile	645	650	655	
Phe	Val	Glu	Ser	Glu	Leu	Pro	Leu	Ser	Trp	Ala	Asp	Arg	Leu	Ser	Gly	660	665	670	
Cys	Leu	Arg	Gly	Pro	Trp	Ala	Trp	Leu	Val	Val	Leu	Leu	Ala	Met	Leu	675	680	685	
Val	Glu	Val	Ala	Leu	Cys	Thr	Trp	Tyr	Leu	Val	Ala	Phe	Pro	Pro	Glu	690	695	700	
Val	Val	Thr	Asp	Trp	His	Met	Leu	Pro	Thr	Glu	Ala	Leu	Val	His	Cys	705	710	715	720
Arg	Thr	Arg	Ser	Trp	Val	Ser	Phe	Gly	Leu	Ala	His	Ala	Thr	Asn	Ala	725	730	735	
Thr	Leu	Ala	Phe	Leu	Cys	Phe	Leu	Gly	Thr	Phe	Leu	Val	Arg	Ser	Gln	740	745	750	
Pro	Gly	Cys	Tyr	Asn	Arg	Ala	Arg	Gly	Leu	Thr	Phe	Ala	Met	Leu	Ala	755	760	765	
Tyr	Phe	Ile	Thr	Trp	Val	Ser	Phe	Val	Pro	Leu	Leu	Ala	Asn	Val	Gln	770	775	780	
Val	Val	Leu	Arg	Pro	Ala	Val	Gln	Met	Gly	Ala	Leu	Leu	Leu	Cys	Val	785	790	795	800
Leu	Gly	Ile	Leu	Ala	Ala	Phe	His	Leu	Pro	Arg	Cys	Tyr	Leu	Leu	Met	805	810	815	
Arg	Gln	Pro	Gly	Leu	Asn	Thr	Pro	Glu	Phe	Phe	Leu	Gly	Gly	Gly	Pro	820	825	830	
Gly	Asp	Ala	Gln	Gly	Gln	Asn	Asp	Gly	Asn	Thr	Gly	Asn	Gln	Gly	Lys	835	840	845	
His	Glu															850			

<210> 5
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> modified_base
 <222> (3)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base

<222> (9)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base
 <222> (12)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base
 <222> (18)
 <223> a, c, t, g, other or unknown

<220>
 <223> Description of Artificial Sequence: Primer

<400> 5
 cgnttyytng cntggggnga rcc

23

<210> 6
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> modified_base
 <222> (3)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base
 <222> (6)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base
 <222> (18)
 <223> a, c, t, g, other or unknown

<220>
 <221> modified_base
 <222> (21)
 <223> a, c, t, g, other or unknown

<220>
 <223> Description of Artificial Sequence: Primer

<400> 6
 cgngcncgrt trtarcanc ngg

23

<210> 7
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 7
Arg Phe Leu Ala Trp Gly Glu Pro Ala
1 5

```
<210> 8
<211> 8
<212> PRT
<213> Homo sapiens
```

```
<400> 8
Pro Gly Cys Tyr Asn Arg Ala Arg
  1             5
```

<210> 9
<211> 552
<212> DNA
<213> Mus sp.

<400> 9						
gtgctgtcac	tcctcctgct	gctttgcctg	gtgctgggtc	tagcactggc	tgctctgggg	60
ctctctgtcc	accactggga	cagccctctt	gtccaggcct	caggcggttc	acagtctctg	120
tttggcctga	tctgcctagg	cctcttctgc	ctcagtgctc	ttctgttccc	aggacggcca	180
agctctgcc	gctgccttgc	acaacaacca	atggctcacc	tcctctcac	aggctgcctg	240
agcacactct	tcctgcaagc	agctgagacc	tttgtggagt	ctgagctgcc	actgagctgg	300
gaaactggc	tatgcagcta	ccttcgggac	tctggcctgc	tagtggtact	gttggccact	360
tttgtggagg	cagcactatg	tgcttggtat	ttgaccgctt	caccagaagt	ggtgacagac	420
tggtcagtg	tgccacaga	ggtactggag	cactgccacg	tgcgttctctg	ggtcaacctg	480
ggcttggtg	acatcaccaa	tgcaatggta	gcttttctct	gctttctggg	cactttcctg	540
gtacaagacc	ag					552

<210> 10
<211> 184
<212> PRT
<213> Mus sp.

```
<400> 10
Val Leu Ser Leu Leu Leu Leu Cys Leu Val Leu Gly Leu Ala Leu
  1             5             10             15
```

Ala Ala Leu Gly Leu Ser Val His His Trp Asp Ser Pro Leu Val Gln
20 25 30

Ala Ser Gly Gly Ser Gln Phe Cys Phe Gly Leu Ile Cys Leu Gly Leu
35 40 45

Phe Cys Leu Ser Val Leu Leu Phe Pro Gly Arg Pro Ser Ser Ala Ser
50 55 60

Cys Leu Ala Gln Gln Pro Met Ala His Leu Pro Leu Thr Gly Cys Leu
65 70 75 80

Ser Thr Leu Phe Leu Gln Ala Ala Glu Thr Phe Val Glu Ser Glu Leu
85 90 95

Pro Leu Ser Trp Ala Asn Trp Leu Cys Ser Tyr Leu Arg Asp Ser Gly
 100 105 110

Leu Leu Val Val Leu Leu Ala Thr Phe Val Glu Ala Ala Leu Cys Ala
 115 120 125

Trp Tyr Leu Thr Ala Ser Pro Glu Val Val Thr Asp Trp Ser Val Leu
 130 135 140

Pro Thr Glu Val Leu Glu His Cys His Val Arg Ser Trp Val Asn Leu
 145 150 155 160

Gly Leu Val His Ile Thr Asn Ala Met Val Ala Phe Leu Cys Phe Leu
 165 170 175

Gly Thr Phe Leu Val Gln Asp Gln
 180

<210> 11
 <211> 558
 <212> DNA
 <213> Rattus sp.

<400> 11
 gtgctgtcac ttctcctgct gctttgcctg gtgctgggccc tgacactggc tgccttgggg 60
 ctctttgtcc actactggga cagccctctt gttcaggcct caggtggggtc actgttctgc 120
 tttggcctga tctgcctagg cctcttctgc ctgagtgcc ttctgttccc aggacgacca 180
 cgctctgcca gctgccttgc ccaacaacca atggctcacc tccctctcac aggctgcctg 240
 agcacactct tcttgcgaagc agccgagatc tttgtggagt ctgagctgcc actgagttgg 300
 gcaaactggc tctgcagcta ccttcggggc ccctgggctt ggctgggtggg actgctggcc 360
 actcttgtgg aggtgcact atgtgcctgg tacttgatgg ctttccctcc agaggtgggtg 420
 acagattggc aggtgctgcc cacggaggtg ctggaacact gccgcatgcg ttcctgggtc 480
 agcctgggct tgggtgcacat caccaatgca ggggtagctt tctctgctt tctgggcact 540
 ttcttggtac aaagccag 558

<210> 12
 <211> 186
 <212> PRT
 <213> Rattus sp.

<400> 12
 Val Leu Ser Leu Leu Leu Leu Cys Leu Val Leu Gly Leu Thr Leu
 1 5 10 15
 Ala Ala Leu Gly Leu Phe Val His Tyr Trp Asp Ser Pro Leu Val Gln
 20 25 30
 Ala Ser Gly Gly Ser Leu Phe Cys Phe Gly Leu Ile Cys Leu Gly Leu
 35 40 45
 Phe Cys Leu Ser Val Leu Leu Phe Pro Gly Arg Pro Arg Ser Ala Ser
 50 55 60
 Cys Leu Ala Gln Gln Pro Met Ala His Leu Pro Leu Thr Gly Cys Leu
 65 70 75 80

Ser Thr Leu Phe Leu Gln Ala Ala Glu Ile Phe Val Glu Ser Glu Leu
 85 90 95
 Pro Leu Ser Trp Ala Asn Trp Leu Cys Ser Tyr Leu Arg Gly Pro Trp
 100 105 110
 Ala Trp Leu Val Val Leu Leu Ala Thr Leu Val Glu Ala Ala Leu Cys
 115 120 125
 Ala Trp Tyr Leu Met Ala Phe Pro Pro Glu Val Val Thr Asp Trp Gln
 130 135 140
 Val Leu Pro Thr Glu Val Leu Glu His Cys Arg Met Arg Ser Trp Val
 145 150 155 160
 Ser Leu Gly Leu Val His Ile Thr Asn Ala Gly Val Ala Phe Leu Cys
 165 170 175
 Phe Leu Gly Thr Phe Leu Val Gln Ser Gln
 180 185

<210> 13
 <211> 2577
 <212> DNA
 <213> Rattus sp.

<400> 13
 atgccgggtt tggctatctt gggcctcagt ctggctgctt tccctggagct tgggatgggg 60
 tcctctttgt gtctgtcaca gcaattcaag gcacaagggg actatatatt ggggtggacta 120
 tttcccctgg gcacaactga ggaggccact ctcaaccaga gaacacagcc caacggcatc 180
 ctatgtacca ggttctcgcc ccttggtttg ttcctggcca tggctatgaa gatggctgta 240
 gaggagatca acaatggatc tgccttgctc cctgggctgc gactgggcta tgacctgttt 300
 gacacatgct cagagccagt ggtcaccatg aagcccagcc tcatgttcat ggccaaggtg 360
 ggaagtcaaa gcattgctgc ctactgcaac tacacacagt accaaccocg tgtgctggct 420
 gtcattgggc cccactcatc agagcttgcc ctctattacag gcaagttctt cagcttcttc 480
 ctcatgccac aggtcagcta tagtgccagc atggatcggc taagtgaccg ggaaacattt 540
 ccatccttct tccgcacagt gcccagtgac cgggtgcagc tgcaggccgt tgtgacactg 600
 ttgcagaatt tcagctggaa ctgggtgggt gccttaggta gtgatgatga ctatggccgg 660
 gaaggtctga gcatcttttc tggcttggtc aactcacgag gtatctgcat tgcacacgag 720
 ggccctgggtc cacaacatga cactagtggc caacaattgg gcaaggtggg ggatgtgcta 780
 cgccaagtga accaaagcaa agtacagggt gtgggtgctgt ttgcatctgc ccgtgctgtc 840
 tactcccttt ttagctacag catccttcat gacctctcac ccaaggtatg ggtggccagt 900
 gagtccctggc tgacctctga cctgggtcatg acacttccca atattgcccg tgtgggact 960
 gttcttgggt ttctgcagcg cgggtgcccta ctgcctgaat tttccatta tgtggagact 1020
 cgccttgccc tagctgctga cccaacattc tgtgcctccc tgaaagctga gttggatctg 1080
 gaggagcgcg tgatggggcc acgctgttca caatgtgact acatcatgct acagaacctg 1140
 tcatctgggc tgatgcagaa cctatcagct gggcagttgc accaccaaatt atttgcaacc 1200
 tatgcagctg tgtacagtgt ggctcaggcc cttcacaaca cctgcagtg caatgtctca 1260
 cattgccaca catcagagcc tgttcaacct tggcagctcc tggagaacat gtacaatatg 1320
 agtttccgtg ctcgagactt gacactgcag tttgatgcca aagggaggtg agacatggaa 1380
 tatgacctga agatgtgggt gtggcagagc cctacacctg tactacatac tgtaggcacc 1440
 ttcaacggca cccttcagct gcagcactcg aaaatgtatt ggccaggcaa ccaggtgcca 1500
 gtcctccagt gctcccggca gtgcaaatgt gcagaggtgc gcagagtaaa gggctttcat 1560
 tcctgtgtgt atgactgtgt ggactgcaag gcagggagct accggaagca tccagatgac 1620
 ttcacctgta tccatgtgg caaggatcag tgggtcccg aaaaaagcac aacctgctta 1680
 cctgcaggc ccaagtttct ggcttggggg gagccagctg tgctgtcact tctcctgctg 1740
 ctttgccctg tgctgggcct gacactgggt gccctggggc tctttgtcca ctactgggac 1800

```

agccctcttg ttcaggcctc aggtgggtca ctgttctgct ttggcctgat ctgcctagge 1860
ctcttctgcc tcagtgtcct tctgttccca ggacgaccac gctctgccag ctgccttgcc 1920
caacaaccaa tggctcacct cctctcaca ggctgcctga gcacactctt cctgcaagca 1980
gccgagatct ttgtggagtc tgagctgcca ctgagttggg caaactggct ctgcagctac 2040
cttcggggcc cctgggcttg gctggtggtg ctgctggcca ctcttggtga ggctgcacta 2100
tgtgcctggt acttgatggc tttccctcca gaggtgggtga cagattggca ggtgctgccc 2160
acggaggtac tggaacactg ccgcattgct tcctgggtca gcctgggctt ggtgcacatc 2220
accaatgcag tgtagctttt cctctgcttt ctgggcactt tcctggtaca gagccagcct 2280
ggtcgctata accgtgcccg tggcctcacc ttgcctatgc tagcttattt catcatctgg 2340
gtctcttttg tgccctcctt ggctaattgt caggtggcct accagccagc tgtgcagatg 2400
ggtgctatct tattctgtgc cctgggcata ctggccacct tccacctgcc caaatgctat 2460
gtacttctgt ggctgccaga gctcaacacc caggagttct tcctgggaag gagccccaag 2520
gaagcatcag atgggaatag tggtagtagt gaggcaactc ggggacacag tgaatga 2577

```

<210> 14

<211> 858

<212> PRT

<213> Rattus sp.

<400> 14

```

Met Pro Gly Leu Ala Ile Leu Gly Leu Ser Leu Ala Ala Phe Leu Glu
  1                      5                      10                      15

Leu Gly Met Gly Ser Ser Leu Cys Leu Ser Gln Gln Phe Lys Ala Gln
          20                      25                      30

Gly Asp Tyr Ile Leu Gly Gly Leu Phe Pro Leu Gly Thr Thr Glu Glu
          35                      40                      45

Ala Thr Leu Asn Gln Arg Thr Gln Pro Asn Gly Ile Leu Cys Thr Arg
          50                      55                      60

Phe Ser Pro Leu Gly Leu Phe Leu Ala Met Ala Met Lys Met Ala Val
          65                      70                      75                      80

Glu Glu Ile Asn Asn Gly Ser Ala Leu Leu Pro Gly Leu Arg Leu Gly
          85                      90                      95

Tyr Asp Leu Phe Asp Thr Cys Ser Glu Pro Val Val Thr Met Lys Pro
          100                      105                      110

Ser Leu Met Phe Met Ala Lys Val Gly Ser Gln Ser Ile Ala Ala Tyr
          115                      120                      125

Cys Asn Tyr Thr Gln Tyr Gln Pro Arg Val Leu Ala Val Ile Gly Pro
          130                      135                      140

His Ser Ser Glu Leu Ala Leu Ile Thr Gly Lys Phe Phe Ser Phe Phe
          145                      150                      155                      160

Leu Met Pro Gln Val Ser Tyr Ser Ala Ser Met Asp Arg Leu Ser Asp
          165                      170                      175

Arg Glu Thr Phe Pro Ser Phe Phe Arg Thr Val Pro Ser Asp Arg Val
          180                      185                      190

```

Gln Leu Gln Ala Val Val Thr Leu Leu Gln Asn Phe Ser Trp Asn Trp
 195 200 205
 Val Ala Ala Leu Gly Ser Asp Asp Asp Tyr Gly Arg Glu Gly Leu Ser
 210 215 220
 Ile Phe Ser Gly Leu Ala Asn Ser Arg Gly Ile Cys Ile Ala His Glu
 225 230 235 240
 Gly Leu Val Pro Gln His Asp Thr Ser Gly Gln Gln Leu Gly Lys Val
 245 250 255
 Val Asp Val Leu Arg Gln Val Asn Gln Ser Lys Val Gln Val Val Val
 260 265 270
 Leu Phe Ala Ser Ala Arg Ala Val Tyr Ser Leu Phe Ser Tyr Ser Ile
 275 280 285
 Leu His Asp Leu Ser Pro Lys Val Trp Val Ala Ser Glu Ser Trp Leu
 290 295 300
 Thr Ser Asp Leu Val Met Thr Leu Pro Asn Ile Ala Arg Val Gly Thr
 305 310 315 320
 Val Leu Gly Phe Leu Gln Arg Gly Ala Leu Leu Pro Glu Phe Ser His
 325 330 335
 Tyr Val Glu Thr Arg Leu Ala Leu Ala Ala Asp Pro Thr Phe Cys Ala
 340 345 350
 Ser Leu Lys Ala Glu Leu Asp Leu Glu Glu Arg Val Met Gly Pro Arg
 355 360 365
 Cys Ser Gln Cys Asp Tyr Ile Met Leu Gln Asn Leu Ser Ser Gly Leu
 370 375 380
 Met Gln Asn Leu Ser Ala Gly Gln Leu His His Gln Ile Phe Ala Thr
 385 390 395 400
 Tyr Ala Ala Val Tyr Ser Val Ala Gln Ala Leu His Asn Thr Leu Gln
 405 410 415
 Cys Asn Val Ser His Cys His Thr Ser Glu Pro Val Gln Pro Trp Gln
 420 425 430
 Leu Leu Glu Asn Met Tyr Asn Met Ser Phe Arg Ala Arg Asp Leu Thr
 435 440 445
 Leu Gln Phe Asp Ala Lys Gly Ser Val Asp Met Glu Tyr Asp Leu Lys
 450 455 460
 Met Trp Val Trp Gln Ser Pro Thr Pro Val Leu His Thr Val Gly Thr
 465 470 475 480
 Phe Asn Gly Thr Leu Gln Leu Gln His Ser Lys Met Tyr Trp Pro Gly
 485 490 495

Asn	Gln	Val	Pro	Val	Ser	Gln	Cys	Ser	Arg	Gln	Cys	Lys	Asp	Gly	Gln		
			500					505					510				
Val	Arg	Arg	Val	Lys	Gly	Phe	His	Ser	Cys	Cys	Tyr	Asp	Cys	Val	Asp		
			515				520					525					
Cys	Lys	Ala	Gly	Ser	Tyr	Arg	Lys	His	Pro	Asp	Asp	Phe	Thr	Cys	Thr		
			530				535				540						
Pro	Cys	Gly	Lys	Asp	Gln	Trp	Ser	Pro	Glu	Lys	Ser	Thr	Thr	Cys	Leu		
					550					555					560		
Pro	Arg	Arg	Pro	Lys	Phe	Leu	Ala	Trp	Gly	Glu	Pro	Ala	Val	Leu	Ser		
				565					570					575			
Leu	Leu	Leu	Leu	Leu	Cys	Leu	Val	Leu	Gly	Leu	Thr	Leu	Ala	Ala	Leu		
				580				585					590				
Gly	Leu	Phe	Val	His	Tyr	Trp	Asp	Ser	Pro	Leu	Val	Gln	Ala	Ser	Gly		
			595				600					605					
Gly	Ser	Leu	Phe	Cys	Phe	Gly	Leu	Ile	Cys	Leu	Gly	Leu	Phe	Cys	Leu		
			610				615				620						
Ser	Val	Leu	Leu	Phe	Pro	Gly	Arg	Pro	Arg	Ser	Ala	Ser	Cys	Leu	Ala		
					630					635					640		
Gln	Gln	Pro	Met	Ala	His	Leu	Pro	Leu	Thr	Gly	Cys	Leu	Ser	Thr	Leu		
				645					650					655			
Phe	Leu	Gln	Ala	Ala	Glu	Ile	Phe	Val	Glu	Ser	Glu	Leu	Pro	Leu	Ser		
			660					665					670				
Trp	Ala	Asn	Trp	Leu	Cys	Ser	Tyr	Leu	Arg	Gly	Pro	Trp	Ala	Trp	Leu		
			675				680					685					
Val	Val	Leu	Leu	Ala	Thr	Leu	Val	Glu	Ala	Ala	Leu	Cys	Ala	Trp	Tyr		
			690				695				700						
Leu	Met	Ala	Phe	Pro	Pro	Glu	Val	Val	Thr	Asp	Trp	Gln	Val	Leu	Pro		
					710					715				720			
Thr	Glu	Val	Leu	Glu	His	Cys	Arg	Met	Arg	Ser	Trp	Val	Ser	Leu	Gly		
				725					730					735			
Leu	Val	His	Ile	Thr	Asn	Ala	Val	Leu	Ala	Phe	Leu	Cys	Phe	Leu	Gly		
			740					745					750				
Thr	Phe	Leu	Val	Gln	Ser	Gln	Pro	Gly	Arg	Tyr	Asn	Arg	Ala	Arg	Gly		
			755				760					765					
Leu	Thr	Phe	Ala	Met	Leu	Ala	Tyr	Phe	Ile	Ile	Trp	Val	Ser	Phe	Val		
			770				775				780						
Pro	Leu	Leu	Ala	Asn	Val	Gln	Val	Ala	Tyr	Gln	Pro	Ala	Val	Gln	Met		
					790					795				800			

Gly Ala Ile Leu Phe Cys Ala Leu Gly Ile Leu Ala Thr Phe His Leu
805 810 815

Pro Lys Cys Tyr Val Leu Leu Trp Leu Pro Glu Leu Asn Thr Gln Glu
820 825 830

Phe Phe Leu Gly Arg Ser Pro Lys Glu Ala Ser Asp Gly Asn Ser Gly
835 840 845

Ser Ser Glu Ala Thr Arg Gly His Ser Glu
850 855

<210> 15
<211> 8194
<212> DNA
<213> Homo sapiens

<220>
<221> modified_base
<222> (1251)..(1300)
<223> a, c, t, g, other or unknown

<220>
<221> modified base
<222> (1951)..(2000)
<223> a, c, t, g, other or unknown

<400> 15
gagaatctcg cgagatcccg tcggtccgcc ccgctgccct cccagctgcc gaaaagaggg 60
gcctccgagc cgccggcgcc ctctgccggc aacctccgga agcacactag gaggttccag 120
ccgatctggt cgagggggtc cacggaggac tccatttacg ttacgcaaatt tccctacccc 180
agccggccgg agagagaaaag ccagaaaacct cgcgaccagc catggggccac ctctccggaa 240
aaacaccggg atattttttt tctcctgcag aaaaagcttt aggattggca gtttaaacia 300
aacatgtcta tttgcatacc ttcggtttgc atgcatttgt ttcgaagtga gcaaccctgg 360
gtaacaaggc gaaagtatat gacaatttgc tcagaatctt aatgtcagaa aactggagac 420
tggggcaggg ggggtgtcgac tcaaagctgt gtctcattta gtaaactgag gccaggttaa 480
aaagtcttga aacctcgcaa caccgggaga aattgtgttc cagcctccca cctcgcccca 540
aaatgccaga gctccttttc taagccaggt gaagtcacag agcgtggaca gaaccacaa 600
ccgtccagag gaaggggtcac tgggtgccac ctggtttgca tctgtgcctt cgtcctgccc 660
agttcctgag tgggaccgca ggcccggaaat gtcaaggcaa acagtcctgc ttcagccact 720
gggtccagt cccaccctt ttgggggctt gaagtttagga agcatccggc agctgccttc 780
tatttaagca actggcctcc ttagaggcca ctcttggtc atgccaggcg cgggcatctg 840
gccagcatgc tgcctgcac ggctgcctg gtccggcctgc agcttctcat ttctgtgtgc 900
tgggcctttg cctgccatag cagggagtct tctcctgact tcaccctccc cggagattac 960
ctcctggcag gctgtttccc tctccattct ggctgtctgc aggtgaggca cagacccgag 1020
gtgaccctgt gtgacaggtg agtgaggggc cagcagagcc acacttagtg ggacccttg 1080
ctatagggcc cctctggctg ccatectcca aacaggacct tgctctgcc tttgcccctt 1140
gaactgtccc caggccttgt tcatcaatcc acttgccacc taagtgtctg ctagaccttc 1200
ctagacactt cggccagttt ccaattattt cacccttgct gttagaatgt nnnnnnnnnn 1260
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn aattccttaa actaaatttc 1320
tacttttctc tctctctctg gaaaacactg actaatgtag caggtttctc tgctccagga 1380
cttcaggacc ttttcgatgc taataagttt ctccatcagg gccagcttgt tctcctact 1440
gagcttgaga gcccttggtg aagttgtggt ttgggggact ggaccgatga cctcaaaggt 1500
tccctttgct cccaagcctc agagtctagg aggccagagg gtctcagcag gcctttgtcc 1560
ttctcagctg tctcttactg gctttctcca caggctctgt agcttcaatg agcatggcta 1620
ccacctcttc caggctatgc ggcttggggg tgaggagata aacaactcca cggccctgct 1680
gcccacatc accctggggg accagctgta tgatgtgtgt tctgactctg ccaatgtgta 1740

tgccacgctg	agagtgtctt	ccctgccagg	gcaacaccac	atagagctcc	aaggagacct	1800
tctccactat	tcccctacgg	tgttggcagt	gattgggcct	gacagcacca	accgtgtctg	1860
caccacagcc	gccctgtctg	gccctttcct	ggtgcccatg	gtaagctgga	gcctcagacc	1920
tttgcccatc	tcccttcagg	caagtctggg	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	1980
nnnnnnnnnn	nnnnnnnnnn	gccaccatgc	ccggctaatt	tttttgtatt	tttagtagag	2040
acgggggttc	accgtgttag	ccaggctggt	cgcaaactcc	taacctcgtg	atccaccacc	2100
ctcggcctcc	caatgtgtctg	ggattacagg	tgtgagccac	tgcacccggc	cataatgtat	2160
taatataata	aaataattat	acaactcacc	ataatgtaga	atcagtggga	gccctgagct	2220
tgttttccta	caactagatg	gtcccatctg	ggggtgatgg	gagacagtga	cagatcatca	2280
gacattagat	tctcataagt	agcgtgcaac	ccagatccct	cgcagtgtga	gttcacagta	2340
gggttcaagc	tcctacaaga	atctgatgct	gctgctgata	tgacaggagg	ggagcagctg	2400
taaatacaga	tgaagcttcg	cttactcacc	agctgctcac	ctcctcctgt	gaggcccggg	2460
tcctaacagg	ccactgacct	aacttctgcc	ctgacctaca	catgcttctc	ttcttccttg	2520
caaaactgcct	ccagtgggaag	tccctgaagg	tccccaacaa	cacgggacta	tttcaactcct	2580
atgcagggtt	tgtctccttt	gcttggaatg	catccctcca	ccccttgtcc	ccaggcagat	2640
tcccacccct	cccccagaac	ctgccccagt	ggagccttcg	cagggtgatt	gtcagtttca	2700
caggctgagg	ggtgtctctc	tgggtctccc	ggctccctgt	atccccacac	ccagcacagg	2760
gccaggcact	ggggggggcct	tcagtggaga	ctgaatggc	tgaacgggac	ctcccataga	2820
ttagctatgc	ggccagcagc	gagacgctca	gcgtgaagcg	gcagtatccc	tctttcctgc	2880
gcaccatccc	caatgacaag	taccagggtg	agaccatggt	gctgctgctg	cagaagtctg	2940
ggtggacctg	gatctctctg	gttggcagca	gtgacgacta	tgggcagcta	ggggtgcagg	3000
cactggagaa	ccaggccact	ggtcagggga	tctgcattgc	tttcaaggac	atcatgccct	3060
tctctgcccc	ggtgggcgat	gagaggatgc	agtgcctcat	gcgccacctg	gcccaggccg	3120
gggccaccgt	cgtggttggt	ttttccagcc	ggcagttggc	cagggtggtt	ttcgagtccg	3180
tgggtgctgac	caacctgact	ggcaagggtg	gggtgccttc	agaagcctgg	gccctctcca	3240
ggcacatcac	tgggtgcccc	gggatccagc	gcattgggat	ggtgctgggg	gtggccatcc	3300
agaagagggc	gtccctggc	ctgaaggcgt	ttgaagaagc	ctatgcccg	gcagacaaga	3360
aggcccctag	gccttgccac	aagggtcctc	ggtgcagcag	caatcagctc	tgcagagaat	3420
gccaagcttt	catggcacac	acgatgcccc	agctcaaagc	cttctccatg	agttctgcct	3480
acaacgcata	ccgggctgtg	tatgcggtgg	cccatggcct	ccaccagctc	ctgggctgtg	3540
cctctggagc	ttgttccagg	ggccgagctc	accctggcca	ggtaagagag	cccaccccag	3600
cacctcctgt	cagggagaac	agccaatcct	gagatgagca	gagtgggcac	tctccggtca	3660
ctctaaatgc	caagggggat	aaatgccact	aacttgagg	tttttgtttt	gttttgtttt	3720
gtttttttag	acagtctggc	tctgtcaccc	aggctgcagt	gtagtgatgc	gatctcggct	3780
ctctgcaact	tccacctcct	gggttcaagt	gattctcttg	cctcggcctc	ctgagttagt	3840
gggattacag	gcacccacca	ccatgcctgg	ataatttttc	ttttcttttt	tttttttttg	3900
agatagatgc	tcgctctggt	gcccaggctg	gaatgcagtg	gtgcatctt	ggctcactgt	3960
gagctccgcc	tcccagggtt	actccattcc	cctgcctcag	cctcccaagt	agggtgggact	4020
acgggcgccc	gccaccacgc	ccagctaatt	ttttttgtat	tttgagtaga	gacgggggtt	4080
caccatgtta	gccaggatgg	tctcaatctc	ctgaccttgt	catccgcccc	cctcgtcctc	4140
ccaaagtgtc	gggattacag	gcgtgagcca	ccgcacccgg	cctaattttt	gtatttttag	4200
tagagatggg	gtttcaccat	gttggccagg	ctggtctcga	actcctggca	tcaagtgatc	4260
ctcctgcttc	ggcctcccaa	agtgtcggga	ttacaggcat	tagctctctt	ctcttagaca	4320
gatctttctc	tctgatecct	gccttctctc	accactgtg	tcttggaagt	gtcaagtgat	4380
aagatccagg	gctaaaactg	tctgtaaagg	agtgtttggt	agaggcctcc	tctcaggagg	4440
ttggtgggga	agattgaggg	gcttcctaag	aagggaaggga	cgagaccttc	ctgatgggct	4500
gaaaccacca	ggacggaaac	ccaggaaggc	cccaggccct	tgcttctggg	accatgtggg	4560
tctgtgctgt	ctgtggtggc	ttcatgatac	gcgtttcttt	cagcttttgg	agcagatcca	4620
caagggtgat	ttccttctac	acaaggacac	tgtggcggtt	aatgacaaca	gagatccctc	4680
cagtagctat	aacataattg	cctgggactg	gaatggaccc	aagtggacct	tcacgggtcct	4740
cgggttcctcc	acatggtctc	cagttcagct	aaacataaat	gagaccaaaa	tccagtggca	4800
cggaaaggac	aaccaggtaa	tggggatgtg	gctactcacc	atgtaactgg	cttatgggca	4860
acctagagcc	tgggggtgat	gctgacacag	tgtacaggga	gcaggagggg	ggccccaggg	4920
gtccagctgc	caccactcta	cccatcctgg	ccagggaagc	aggggaagaca	ctccgtaggc	4980
gagtggtgag	atgcctggg	gcggaagttc	acacgaccag	gggcccctgc	ctgggagtga	5040
gccctgaggg	cagatgcaca	gagattctgt	tttctgttcc	acatgtgagc	tgctccttga	5100
cttggggccc	tacgtgtggc	ccctctggct	tcttacagg	gcctaagtct	gtgtgttcca	5160
gcgactgtct	tgaagggcac	cagcgagtgg	ttacgggttt	ccatcactgc	tgcttttagt	5220


```

gtgtgccctg tggggctggg accttcctca acaagagtgg tgagtgggca atggagcagg 5280
cgagctaccc agcactcccg ggggctgcac ggtggaggga gggcctccct tgggccccat 5340
gtgccctgcc ccagaaccaa ggcccagtcg ctgggctgcc agttagcttc aggttgaggg 5400
acacctgcta ccagacagaa ttctgatcaa gagaatcagc cactgggtgc ggtggctcat 5460
gcctgtaatc ccagcacttt gggaggctga ggcggtgga tcacttgagg tcgggagttc 5520
gagaccagcc tggccaacat ggtgaaaccc catctctacc aaaaatataa aaaattagct 5580
gggtgtggtg gcgcgtgcct gtaatcccag ctactcggga ggctgaggca ggagaatcac 5640
ttgaacccag gaggcgagg ttgcagtga ccaagatgca ttccagcctg gaccacaaag 5700
cgagaattcg tcccccaaaa aaaagaaagg aggcggggcg cgggtggctca cacctgtaat 5760
cccagcactt tgggaggccg aggtgggtgg atcacctgag gtcaggagtt cgagaccagc 5820
ctgaccaaca tgggtgaaacc ccactctctac taaaaatata aaaaaagtta gccgggcgtt 5880
gtggcgtgtg cctgtaattc cagctactcg ggaggctgag gcaggagaat tgcttgaacc 5940
cgggaggcgg aggttgagct gagccaagat tgcaccattg cactccagcc tgggcgacaa 6000
gagaaaaact ctgtctcaaa aaaaaagaaa gaaagaaaga attagccaac tgaaagcctt 6060
agactgaggt gtgtcctctg ttagagagct gtcatacaaa ctccatacaa agcagtcgta 6120
tcctgaattc aacctctttc tctaaatgaa tatagctatt gttccctttg tgccctcttg 6180
tcctactgtc ccttctgttg cccatgccaa agacagctag ctccctgaac agcttggcct 6240
gaatacagat actagcgtgt ctgcagcaga gaaaaaaaca gcattcccca tccagaaatg 6300
caaggtcaag aacagagagc aaattaggta gctaaggact caggtcctta gttggtgtcc 6360
aggggccaca ttctttcctt tcaccatctc tgtagggaca ggaatacttc cttctgtcc 6420
tcagagggtc aggactcaga gaaaccacag agcagcagct caggaaagtg gttcatggaa 6480
atgctggcaa gagagagggg ttacaatgcc ctcccttggg agcaggctgc tcccatcaga 6540
tcgtaacctc tctggtatgt gggcagagct accaggttaa ggtcctccct agggtttgca 6600
aaaccctcat gggatcatga gccatacaga accgacctgt gtgtctccag agtctgtaat 6660
taacacagcc attttgagga aatgcgtggc ctcaggcccc actcccggt accccatcc 6720
cactatgcct agtatagtct agtgccctg gtacaattct cccagtatct tgcaggcccc 6780
tatttcctat tcctactctg ctcatctggc tctcaggaac cttcttggcc ttccctttca 6840
gacctctaca gatgccagcc ttgtgggaaa gaagagtggg cacctgaggg aagccagacc 6900
tgcttcccgc gcactgtggt gtttttggt ttgcgtgagc acacctcttg ggtgctgctg 6960
gcagctaaca cgctgctgct gctgctgctg cttgggactg ctggcctgtt tgectggcac 7020
ctagacaccc ctgtggtgag gtcagcaggg ggccgcctgt gctttcttat gctgggctcc 7080
ctggcagcag gtagtggcag cctctatggc ttctttgggg aaccacaaag gcctgcgtgc 7140
ttgctacgcc aggcctctt tgcccttggg ttccaccatc tcctgtcctg cctgacagtt 7200
cgctcattcc aactaatcat catcttcaag ttttcacca aggtacctac attctaccac 7260
gcctgggtcc aaaaccacgg tgctggcctg tttgtgatga tcagctcagc ggcccagctg 7320
cttatctgtc taacttggct ggtggtgtgg accccactgc ctgctaggga ataccagcgc 7380
ttcccccatc tgggtgatgt tgagtgcaca gagaccaact ccctgggctt catactggcc 7440
ttcctctaca atggcctcct ctccatcagt gcctttgcct gcagctacct gggtaaggac 7500
ttgccagaga actacaacga ggccaaatgt gtcaccttca gcctgctctt caacttcgtg 7560
tcctggatcg ccttcttcac cacggccagc gtctacgacg gcaagtacct gcctgcggcc 7620
aacatgatgg ctgggctgag cagcctgagc agcggcttcg gtgggtattt tctgcctaag 7680
tgctacgtga tcctctgccg cccagacctc aacagcacag agcacttcca ggcctccatt 7740
caggactaca cgaggcgtg cggctccacc tgaccagtgg gtcagcaggc acggctggca 7800
gccttctctg ccctgagggt cgaaggctga gcaggccggg ggtgtccggg aggtctttgg 7860
gcatcgcggt ctgggggttg gacgtgtaag cgccctggag agcctagacc aggtccggg 7920
ctgccaataa agaagtgaag tgcgtatctg gtctcctgtc gtgggagagt gtgaggtgta 7980
acggattcaa gtctgaaccc agagcctgga aaaggctgac cgcccagatt gacgttgcta 8040
ggcaactccg gaggcgggcc cagcgccaaa agaacagggc gaggcgtcgt ccccgcatcc 8100
cattggccgt tctctgcggg gcccgccct cggggggccg agctagaagc tctacgttc 8160
cgaggcgcac ctccctggcct gcacgctttg acgt 8194

```

<210> 16

<211> 2526

<212> DNA

<213> Homo sapiens

<400> 16

```

atgctgctct gcacgggctcg cctgggtcggc ctgcagcttc tcatttctctg ctgctggggcc 60
tttgccctgcc atagcacgga gtcttctcct gacttcaccc tccccggaga ttacctcctg 120
gcaggcctgt tccctctcca ttctggctgt ctgcaggtga ggcacagacc cgaggtgacc 180
ctgtgtgaca ggtctttag tagctatgag catgggtacc acctcttcca ggctatgcgg 240
cttgggggttg aggagataaa caactccacg gccctgctgc ccaacatcac cctgggggtac 300
cagctgtatg atgtgtgttc tgactctgcc aatgtgtatg ccacgctgag agtgcctccc 360
ctgccagggc aacaccacat agagctccaa ggagaccttc tccactattc ccctacgggtg 420
ctggcagtga ttgggcctga cagcaccaac cgtgctgcca ccacagccgc cctgctgagc 480
cctttcctgg tgcccatgat tagctatgag gccagcagcg agacgctcag cgtgaagcgg 540
cagtatccct ctttctctgcg caccatcccc aatgacaagt accagggtgga gaccatgggtg 600
ctgctgctgc agaagttcgg gtggacctgg atctctctgg ttggcagcag tgacgactat 660
gggcagctag ggggtgcaggc actggagaac caggccactg gtcaggggat ctgcattgct 720
ttcaaggaca tcatgccctt ctctgcccag gtgggcgatg agaggatgca gtgcctcatg 780
cgccacctgg cccaggccgg ggccaccgtc gtggttggtt tttccagccg gcagttggcc 840
aggggtgttt tcgagtcctg ggtgctgacc aacctgactg gcaagggtgtg ggtcgcctca 900
gaagcctggg cctctctccag gcacatcact ggggtgcccc ggatccagcg cattgggatg 960
gtgctggggc tggccatcca gaagagggtg gtccttgccc tgaaggcgtt tgaagaagcc 1020
tatgcccggg cagacaagaa ggcccctagg ccttgccaca agggctcctg gtgcagcagc 1080
aatcagctct gcagagaatg ccaagctttc atggcacaca cgatgccccaa gctcaaagcc 1140
ttctccatga gttctgcta caacgcatac cgggctgtgt atgcgggtggc ccatggcctc 1200
caccagctcc tgggctgtgc ctctggagct tgttccaggg gccgagtcta cccctggcag 1260
cttttgagc agatccacaa ggtgcatttc cttctacaca aggacactgt ggcgtttaat 1320
gacaacagag atccccctag tagctataac ataattgcct gggactggaa tggacccaag 1380
tggaccttca cggctcctcg ttcctccaca tgggtctccag ttcagctaaa cataaatgag 1440
acaaaaatcc agtggcacgg aaaggacaac cagggtgccta agtctgtgtg ttccagcgac 1500
tgtcttgaag ggcaccagcg agtggttacg ggtttccatc actgctgctt tgagtgtgtg 1560
ccctgtgggg ctgggacctt cctcaacaag agtgacctct acagatgcca gccttgtggg 1620
aaagaagagt gggcacctga ggggaagccag acctgcttcc cgcgcactgt ggtgtttttg 1680
gctttgcgtg agcacacctc ttgggtgctg ctggcagcta acacgctgct gctgctgctg 1740
ctgcttggga ctgctggcct gtttgctggg cactagaca cccctgtggg gaggtcagca 1800
gggggcccgc tgtgctttct tatgctgggc tccctggcag caggtagtgg cagcctctat 1860
ggcttctttg ggggaaccac aaggcctgcg tgcttgctac gccaggccct ctttgccctt 1920
ggtttcacca tcttctctgc ctgcctgaca gttcgctcat tccaaactaat catcatcttc 1980
aagttttcca ccaaggtacc tacattctac cagccttggg tccaaaacca cgggtgctggc 2040
ctggtttgta tgatcagctc agcggccag ctgcttatct gtctaacttg gctgggtggg 2100
tggaaccac tgacctag ggaataccag cgcttcccc atctgggtgat gcttgagtgc 2160
acagagacca actccctggg cttcactact gccttctctt acaatggcct cctctccatc 2220
agtgcctttg cctgcagcta cctgggtgag gacttgccag agaactacaa cgaggccaaa 2280
tgtgtcacct tcagcctgct cttcaacttc gtgtcctgga tcgccttctt caccacggcc 2340
agcgtctacg acggcaagta cctgcctgcg gccaacatga tggctgggct gagcagcctg 2400
agcagcggct tcggtgggta ttttctgcct aagtgtacg tgatcctctg ccgccagac 2460
ctcaacagca cagagcactt ccaggcctcc attcaggact acacgaggcg ctgcggctcc 2520
acctga

```

<210> 17

<211> 841

<212> PRT

<213> Homo sapiens

<400> 17

```

Met Leu Leu Cys Thr Ala Arg Leu Val Gly Leu Gln Leu Leu Ile Ser
  1             5             10             15

```

```

Cys Cys Trp Ala Phe Ala Cys His Ser Thr Glu Ser Ser Pro Asp Phe
  20             25             30

```

Thr Leu Pro Gly Asp Tyr Leu Leu Ala Gly Leu Phe Pro Leu His Ser
 35 40 45
 Gly Cys Leu Gln Val Arg His Arg Pro Glu Val Thr Leu Cys Asp Arg
 50 55 60
 Ser Cys Ser Phe Asn Glu His Gly Tyr His Leu Phe Gln Ala Met Arg
 65 70 75 80
 Leu Gly Val Glu Glu Ile Asn Asn Ser Thr Ala Leu Leu Pro Asn Ile
 85 90 95
 Thr Leu Gly Tyr Gln Leu Tyr Asp Val Cys Ser Asp Ser Ala Asn Val
 100 105 110
 Tyr Ala Thr Leu Arg Val Leu Ser Leu Pro Gly Gln His His Ile Glu
 115 120 125
 Leu Gln Gly Asp Leu Leu His Tyr Ser Pro Thr Val Leu Ala Val Ile
 130 135 140
 Gly Pro Asp Ser Thr Asn Arg Ala Ala Thr Thr Ala Ala Leu Leu Ser
 145 150 155 160
 Pro Phe Leu Val Pro Met Ile Ser Tyr Ala Ala Ser Ser Glu Thr Leu
 165 170 175
 Ser Val Lys Arg Gln Tyr Pro Ser Phe Leu Arg Thr Ile Pro Asn Asp
 180 185 190
 Lys Tyr Gln Val Glu Thr Met Val Leu Leu Leu Gln Lys Phe Gly Trp
 195 200 205
 Thr Trp Ile Ser Leu Val Gly Ser Ser Asp Asp Tyr Gly Gln Leu Gly
 210 215 220
 Val Gln Ala Leu Glu Asn Gln Ala Thr Gly Gln Gly Ile Cys Ile Ala
 225 230 235 240
 Phe Lys Asp Ile Met Pro Phe Ser Ala Gln Val Gly Asp Glu Arg Met
 245 250 255
 Gln Cys Leu Met Arg His Leu Ala Gln Ala Gly Ala Thr Val Val Val
 260 265 270
 Val Phe Ser Ser Arg Gln Leu Ala Arg Val Phe Phe Glu Ser Val Val
 275 280 285
 Leu Thr Asn Leu Thr Gly Lys Val Trp Val Ala Ser Glu Ala Trp Ala
 290 295 300
 Leu Ser Arg His Ile Thr Gly Val Pro Gly Ile Gln Arg Ile Gly Met
 305 310 315 320
 Val Leu Gly Val Ala Ile Gln Lys Arg Ala Val Pro Gly Leu Lys Ala
 325 330 335

Phe Glu Glu Ala Tyr Ala Arg Ala Asp Lys Lys Ala Pro Arg Pro Cys
 340 345 350
 His Lys Gly Ser Trp Cys Ser Ser Asn Gln Leu Cys Arg Glu Cys Gln
 355 360 365
 Ala Phe Met Ala His Thr Met Pro Lys Leu Lys Ala Phe Ser Met Ser
 370 375 380
 Ser Ala Tyr Asn Ala Tyr Arg Ala Val Tyr Ala Val Ala His Gly Leu
 385 390 395 400
 His Gln Leu Leu Gly Cys Ala Ser Gly Ala Cys Ser Arg Gly Arg Val
 405 410 415
 Tyr Pro Trp Gln Leu Leu Glu Gln Ile His Lys Val His Phe Leu Leu
 420 425 430
 His Lys Asp Thr Val Ala Phe Asn Asp Asn Arg Asp Pro Leu Ser Ser
 435 440 445
 Tyr Asn Ile Ile Ala Trp Asp Trp Asn Gly Pro Lys Trp Thr Phe Thr
 450 455 460
 Val Leu Gly Ser Ser Thr Trp Ser Pro Val Gln Leu Asn Ile Asn Glu
 465 470 475 480
 Thr Lys Ile Gln Trp His Gly Lys Asp Asn Gln Val Pro Lys Ser Val
 485 490 495
 Cys Ser Ser Asp Cys Leu Glu Gly His Gln Arg Val Val Thr Gly Phe
 500 505 510
 His His Cys Cys Phe Glu Cys Val Pro Cys Gly Ala Gly Thr Phe Leu
 515 520 525
 Asn Lys Ser Asp Leu Tyr Arg Cys Gln Pro Cys Gly Lys Glu Glu Trp
 530 535 540
 Ala Pro Glu Gly Ser Gln Thr Cys Phe Pro Arg Thr Val Val Phe Leu
 545 550 555 560
 Ala Leu Arg Glu His Thr Ser Trp Val Leu Leu Ala Ala Asn Thr Leu
 565 570 575
 Leu Leu Leu Leu Leu Leu Gly Thr Ala Gly Leu Phe Ala Trp His Leu
 580 585 590
 Asp Thr Pro Val Val Arg Ser Ala Gly Gly Arg Leu Cys Phe Leu Met
 595 600 605
 Leu Gly Ser Leu Ala Ala Gly Ser Gly Ser Leu Tyr Gly Phe Phe Gly
 610 615 620
 Glu Pro Thr Arg Pro Ala Cys Leu Leu Arg Gln Ala Leu Phe Ala Leu
 625 630 635 640

Gly Phe Thr Ile Phe Leu Ser Cys Leu Thr Val Arg Ser Phe Gln Leu
 645 650 655

 Ile Ile Ile Phe Lys Phe Ser Thr Lys Val Pro Thr Phe Tyr His Ala
 660 665 670

 Trp Val Gln Asn His Gly Ala Gly Leu Phe Val Met Ile Ser Ser Ala
 675 680 685

 Ala Gln Leu Leu Ile Cys Leu Thr Trp Leu Val Val Trp Thr Pro Leu
 690 695 700

 Pro Ala Arg Glu Tyr Gln Arg Phe Pro His Leu Val Met Leu Glu Cys
 705 710 715 720

 Thr Glu Thr Asn Ser Leu Gly Phe Ile Leu Ala Phe Leu Tyr Asn Gly
 725 730 735

 Leu Leu Ser Ile Ser Ala Phe Ala Cys Ser Tyr Leu Gly Lys Asp Leu
 740 745 750

 Pro Glu Asn Tyr Asn Glu Ala Lys Cys Val Thr Phe Ser Leu Leu Phe
 755 760 765

 Asn Phe Val Ser Trp Ile Ala Phe Phe Thr Thr Ala Ser Val Tyr Asp
 770 775 780

 Gly Lys Tyr Leu Pro Ala Ala Asn Met Met Ala Gly Leu Ser Ser Leu
 785 790 795 800

 Ser Ser Gly Phe Gly Gly Tyr Phe Leu Pro Lys Cys Tyr Val Ile Leu
 805 810 815

 Cys Arg Pro Asp Leu Asn Ser Thr Glu His Phe Gln Ala Ser Ile Gln
 820 825 830

 Asp Tyr Thr Arg Arg Cys Gly Ser Thr
 835 840

<210> 18
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MOD_RES
 <222> (1)
 <223> Thr or Arg

<220>
 <221> MOD_RES
 <222> (3)
 <223> Phe or Leu

<220>
 <221> MOD_RES

```

<220>
<221> MOD_RES
<222> (3)
<223> Glu, Gly or Thr

```

<220>
 <221> MOD_RES
 <222> (4)
 <223> Asn, Arg or Cys

<220>
 <221> MOD_RES
 <222> (7)
 <223> Arg or Glu

<220>
 <221> MOD_RES
 <222> (9)
 <223> Arg or Lys

<220>
 <221> MOD_RES
 <222> (10)
 <223> Cys, Gly or Phe

<220>
 <221> MOD_RES
 <222> (11)
 <223> Val, Leu or Ile

<220>
 <221> MOD_RES
 <222> (13)
 <223> Phe or Leu

<220>
 <221> MOD_RES
 <222> (14)
 <223> Ala or Asn

<220>
 <221> MOD_RES
 <222> (15)
 <223> Met or Leu

<220>
 <223> Description of Artificial Sequence: Consensus
 sequence

<400> 19
 Xaa Pro Xaa Xaa Tyr Asn Xaa Ala Xaa Xaa Xaa Thr Xaa Xaa Xaa
 1 5 10 15

<210> 20
 <211> 3563
 <212> DNA
 <213> Homo sapiens

<400> 20
 agcctggcag tggcctcagg cagagtctga cgcgacacaaa ctttcaggcc caggaagcga 60
 ggacaccact ggggccccag ggtgtggcaa gtgaggatgg caagggtttt gctaaacaaa 120
 tcctctgccc gtcctccgcc ccgggctcac tccatgtgag gccccagtcg gggcagccac 180

ctgccgtgcc	tgttggaagt	tgccctctgcc	atgtctggggc	ctgctgtcct	gggcctcagc	240
ctctggggctc	tcctgcaccc	tgggacgggg	gccccattgt	gcctgtcaca	gcaacttagg	300
atgaaggggg	actacgtgct	gggggggctg	ttccccctgg	gcgaggccga	ggaggctggc	360
ctccgcagcc	ggacacggcc	cagcagccct	gtgtgcacca	ggtacagagg	tgggacggcc	420
tgggtcgggg	tcagggtgac	caggtctggg	gtgctcctga	gctggggccg	aggtggccat	480
ctgcggttct	gtgtggcccc	aggttctcct	caaacggcct	gctctgggca	ctggccatga	540
aaatggccgt	ggaggagatc	aacaacaagt	cggatctgct	gccccgggctg	cgccctgggt	600
acgacctctt	tgatacgtgc	tcggagccctg	tgggtggccat	gaagcccagc	ctcatgttcc	660
tggccaaggc	aggcagccgc	gacatcgccg	cctactgcaa	ctacacgcag	taccagcccc	720
gtgtgctggc	tgatcatggg	ccccactcgt	cagagctcgc	catggtcacc	ggcaagttct	780
tcagcttctt	cctcatgccc	cagtggggcg	ccccccacca	tcacccaccc	ccaaccaacc	840
cctgccccgt	gggagccccct	tgtgtcagga	gaatgctaca	tgcacccccc	ccagccctgc	900
cctgggagcc	ctgtgtcaga	agatgctctt	ggccttgtag	gtcagctacg	gtgctagcat	960
ggagctgctg	agcgcgccgg	agaccttccc	ctccttcttc	cgcaccgtgc	ccagcgaccg	1020
tgtgcagctg	acggccgcgg	cggagctgct	gcaggagtgc	ggctggaact	gggtggccgc	1080
cctgggcagc	gacgacgagt	acggccggca	gggcctgagc	atcttctcgg	ccctggccgc	1140
ggcacgcggc	atctgcatcg	cgcacgaggg	cctgggtgccg	ctgccccgtg	ccgatgactc	1200
gcggctgggg	aaggtgcagg	acgtcctgca	ccaggtgaac	cagagcagcg	tgcaggtggt	1260
gctgtctgtc	gcctccgtgc	acgcgcgcca	cgccctcttc	aactacagca	tcagcagcag	1320
gctctcgcgc	aaggtgtggg	tggccagcga	ggcctggctg	acctctgacc	tggtcatggg	1380
gctgccccgc	atggcccaga	tgggcacggg	gcttggtctc	ctccagaggg	gtgcccagct	1440
gcacgagttc	ccccagtagc	tgaagacgca	cctggccctg	gccaccgacc	cggccttctg	1500
ctctgccctg	ggcgagaggg	agcagggctc	ggaggaggac	gtgggtgggg	agcgctgccc	1560
gcagtgtgac	tgcatcacgc	tgcagaacgt	gagcgcaggg	ctaaatcacc	accagacggt	1620
ctctgtctac	gcagctgtgt	atagcgtggc	ccaggccctg	cacaacactc	ttcagtgcaa	1680
cgcctcaggg	tgccccgcgc	aggacccctg	gaagccctgg	caggtgagcc	cgggagatgg	1740
gggtgtgctg	tcctctgcat	gtgcccaggg	caccaggcac	ggccaccacg	cctgacttgg	1800
aggtggctgg	cggtcagcc	cgtcccccg	cccgagctc	ctggagaaca	tgtacaacct	1860
gaccttccac	gtgggcgggc	tgccgctgcg	gttcgacagc	agcggaaacg	tggacatgga	1920
gtacgacctg	aagctgtggg	tgtggcaggg	ctcagtgcgc	aggctccacg	acgtgggcag	1980
gttcaacggc	agcctcagga	cagagcgcct	gaagatccgc	tggcacacgt	ctgacaacca	2040
gggtgaggtga	gggtgggtgt	gccaggcggt	cccgtggtag	cccccgcggc	agggcgagc	2100
ctgggggtgg	gggcccgttc	agtctcccgt	gggcagtgcc	agccgagcag	agccagaccc	2160
caggcctgtg	cgcagaagcc	cgtgtcccgg	tgctcgcggc	agtgccagga	gggccaggtg	2220
cgcggggtca	aggggttcca	ctcctgctgc	tacgactgtg	tggactgcga	ggcgggcagc	2280
taccggcaaa	accaggtga	gcgccttccc	cggcaggcgg	gggtgggaac	gcagcagggg	2340
aggttctctg	caagtcctga	ctctgagacc	agagcccaca	gggtacaaga	cgaacaccca	2400
gcgcccctct	cctctctcac	agacgacatc	gcctgcacct	tttgtggcca	ggatgagttg	2460
tccccggagc	gaagcacacg	ctgcttccgc	cgcaggtctc	ggttcctggc	atggggcgag	2520
ccggctgtgc	tgtgtgtgct	cctgctgctg	agcctggcgc	tgggccttgt	gctggctgct	2580
ttggggctgt	tcgttcacca	tcgggacagc	ccactggttc	aggcctcggg	ggggccccctg	2640
gcctgctttg	gcctgggtgtg	cctgggcctg	gtctgcctca	gcgtcctcct	gttccctggc	2700
cagcccagcc	ctgcccgatg	cctggccccg	cagcccttgt	cccacctccc	gtcacggggc	2760
tgcttgagca	cactcttctc	gcaggcgggc	gagatcttcg	tggagtcaga	actgcctctg	2820
agctgggcag	accggctgag	tggctgctgt	cgggggcccc	gggctgggt	ggtggtgctg	2880
ctggccatgc	tgggtgaggt	cgcactgtgc	acctggtacc	tgggtggcctt	cccgccggag	2940
gtgggtgacgg	actggcacat	gctgcccacg	gaggcgctgg	tgactgccc	cacacgctcc	3000
tgggtcagct	tcggccatgc	gcacgccacc	aatgccacgc	tggcctttct	ctgcttctctg	3060
ggcactttcc	tgggtgcggag	ccagccgggc	tgctacaacc	gtgcccggtg	cctcaccttt	3120
gccatgctgg	cctacttcat	cacctgggtc	tcctttgtgc	ccctcctggc	caatgtgcag	3180
gtggtcctca	ggcccgcctg	gcagatgggc	gccctcctgc	tctgtgtcct	gggcatcctg	3240
gctgccttcc	acctgcccag	gtgttacctg	ctcatgcggc	agccagggct	caacaccccc	3300
gagttcttcc	tgggaggggg	ccctggggat	gccccaggcc	agaatgacgg	gaacacagga	3360
aatcagggga	aacatgagtg	acccaacctt	gtgatctcag	ccccggtgaa	cccagactta	3420
gctgcgatcc	cccccaagcc	agcaatgacc	cgtgtctcgc	tacagagacc	ctcccgctct	3480
aggttctgac	cccaggttgt	ctcctgacct	tgacccca	gtgagcccta	ggcctggagc	3540
acgtggacac	ccctgtgacc	atc				3563